

#8



PCT09

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/936,205

DATE: 02/08/2002
TIME: 11:26:41

Input Set : A:\37945-0024.txt
Output Set: N:\CRF3\02082002\I936205.raw

ENTERED

4 <110> APPLICANT: SMITH, Richard Anthony Godwin
5 PRATT, Julian Roy
6 SACKS, Steven Howard
8 <120> TITLE OF INVENTION: ORGAN TRANSPLANT SOLUTIONS CONTAINING CONJUGATES OF
9 SOLUBLE PEPTIDIC COMPOUNDS WITH MEMBRANE-BINDING
11 <130> FILE REFERENCE: 37945-0024
13 <140> CURRENT APPLICATION NUMBER: US 09/936,205
C--> 14 <141> CURRENT FILING DATE: 2001-12-07
16 <150> PRIOR APPLICATION NUMBER: PCT/GB00/00834
17 <151> PRIOR FILING DATE: 2000-03-08
19 <150> PRIOR APPLICATION NUMBER: GB 9905503.0
20 <151> PRIOR FILING DATE: 1999-03-10
22 <160> NUMBER OF SEQ ID NOS: 11
24 <170> SOFTWARE: PatentIn Ver. 2.1
26 <210> SEQ ID NO: 1
27 <211> LENGTH: 215
28 <212> TYPE: PRT
29 <213> ORGANISM: Artificial Sequence
31 <220> FEATURE:
32 <223> OTHER INFORMATION: Linear, 2 polypeptide chains disulphide linked
34 <220> FEATURE:
35 <221> NAME/KEY: DISULFID
36 <222> LOCATION: (198)..(199)
38 <220> FEATURE:
39 <223> OTHER INFORMATION: 2nd polypeptide chain (199-215) runs C to N
40 terminus
42 <220> FEATURE:
43 <223> OTHER INFORMATION: An N-myristoyl group is at the N-terminus of the
44 second polypeptide chain
46 <220> FEATURE:
47 <223> OTHER INFORMATION: A CONH2 group is at the C terminus of the second
48 polypeptide chain
50 <220> FEATURE:
51 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
52 peptide reagent
54 <400> SEQUENCE: 1
55 Met Gln Cys Asn Ala Pro Glu Trp Leu Pro Phe Ala Arg Pro Thr Asn
56 1 5 10 15
58 Leu Thr Asp Glu Phe Glu Phe Pro Ile Gly Thr Tyr Leu Asn Tyr Glu
59 20 25 30
61 Cys Arg Pro Gly Tyr Ser Gly Arg Pro Phe Ser Ile Ile Cys Leu Lys
62 35 40 45
64 Asn Ser Val Trp Thr Gly Ala Lys Asp Arg Cys Arg Arg Lys Ser Cys

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65      50      55      60
67 Arg Asn Pro Pro Asp Pro Val Asn Gly Met Val His Val Ile Lys Gly
68 65      70      75      80
70 Ile Gln Phe Gly Ser Gln Ile Lys Tyr Ser Cys Thr Lys Gly Tyr Arg
71      85      90      95
73 Leu Ile Gly Ser Ser Ser Ala Thr Cys Ile Ile Ser Gly Asp Thr Val
74      100      105      110
76 Ile Trp Asp Asn Glu Thr Pro Ile Cys Asp Arg Ile Pro Cys Gly Leu
77      115      120      125
79 Pro Pro Thr Ile Thr Asn Gly Asp Phe Ile Ser Thr Asn Arg Glu Asn
80      130      135      140
82 Phe His Tyr Gly Ser Val Val Thr Tyr Arg Cys Asn Pro Gly Ser Gly
83 145      150      155      160
85 Gly Arg Lys Val Phe Glu Leu Val Gly Glu Pro Ser Ile Tyr Cys Thr
86      165      170      175
88 Ser Asn Asp Asp Gln Val Gly Ile Trp Ser Gly Pro Ala Pro Gln Cys
89      180      185      190
91 Ile Ile Pro Asn Lys Cys Cys Asp Gly Pro Lys Lys Lys Lys Lys Lys
92      195      200      205
94 Ser Pro Ser Lys Ser Ser Gly
95      210      215
98 <210> SEQ ID NO: 2
99 <211> LENGTH: 218
100 <212> TYPE: PRT
101 <213> ORGANISM: Artificial Sequence
103 <220> FEATURE:
104 <223> OTHER INFORMATION: 2 polypeptide chains disulphide linked
106 <220> FEATURE:
107 <221> NAME/KEY: DISULFID
108 <222> LOCATION: (198)..(199)
110 <220> FEATURE:
111 <223> OTHER INFORMATION: The second polypeptide chain (199-218) runs C to N
112 terminus
114 <220> FEATURE:
115 <223> OTHER INFORMATION: An N-Myristoyl group is at the N terminus of the
116 second polypeptide chain
118 <220> FEATURE:
119 <223> OTHER INFORMATION: A CONH2 group is at the C terminus of the second
120 polypeptide chain
122 <220> FEATURE:
123 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
124 peptide reagent
126 <400> SEQUENCE: 2
127 Met Gln Cys Asn Ala Pro Glu Trp Leu Pro Phe Ala Arg Pro Thr Asn
128 1      5      10      15
130 Leu Thr Asp Glu Phe Glu Phe Pro Ile Gly Thr Tyr Leu Asn Tyr Glu
131      20      25      30
133 Cys Arg Pro Gly Tyr Ser Gly Arg Pro Phe Ser Ile Ile Cys Leu Lys
134      35      40      45

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136 Asn Ser Val Trp Thr Gly Ala Lys Asp Arg Cys Arg Arg Lys Ser Cys
137 50 55 60
139 Arg Asn Pro Pro Asp Pro Val Asn Gly Met Val His Val Ile Lys Gly
140 65 70 75 80
142 Ile Gln Phe Gly Ser Gln Ile Lys Tyr Ser Cys Thr Lys Gly Tyr Arg
143 85 90 95
145 Leu Ile Gly Ser Ser Ser Ala Thr Cys Ile Ile Ser Gly Asp Thr Val
146 100 105 110
148 Ile Trp Asp Asn Glu Thr Pro Ile Cys Asp Arg Ile Pro Cys Gly Leu
149 115 120 125
151 Pro Pro Thr Ile Thr Asn Gly Asp Phe Ile Ser Thr Asn Arg Glu Asn
152 130 135 140
154 Phe His Tyr Gly Ser Val Thr Tyr Arg Cys Asn Pro Gly Ser Gly
155 145 150 155 160
157 Gly Arg Lys Val Phe Glu Leu Val Gly Glu Pro Ser Ile Tyr Cys Thr
158 165 170 175
160 Ser Asn Asp Asp Gln Val Gly Ile Trp Ser Gly Pro Ala Pro Gln Cys
161 180 185 190
163 Ile Ile Pro Asn Lys Cys Cys Ala Asp Leu Arg Ser Ser Leu Gly Pro
164 195 200 205
166 Lys Lys Lys Lys Lys Lys Ser Pro Ser Gly
167 210 215
170 <210> SEQ ID NO: 3
171 <211> LENGTH: 20
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173 <213> ORGANISM: Artificial Sequence
175 <220> FEATURE:
176 <223> OTHER INFORMATION: An N-myristoyl group is at the N terminus of the
177 polypeptide chain
179 <220> FEATURE:
180 <223> OTHER INFORMATION: A CONH2 group is at the C-terminus of the
181 polypeptide chain
183 <220> FEATURE:
184 <223> OTHER INFORMATION: An S-2-Thiopyridyl group is attached to the
185 C-terminal cysteine
187 <220> FEATURE:
188 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
189 peptide reagent
191 <400> SEQUENCE: 3
192 Gly Ser Pro Ser Lys Lys Lys Lys Lys Pro Gly Leu Ser Ser Arg
193 1 5 10 15
195 Leu Asp Ala Cys
196 20
199 <210> SEQ ID NO: 4
200 <211> LENGTH: 20
201 <212> TYPE: PRT
202 <213> ORGANISM: Artificial Sequence
204 <220> FEATURE:
205 <223> OTHER INFORMATION: A CONH2 group is at the C terminus

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207 <220> FEATURE:
208 <223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic
209 peptide
211 <400> SEQUENCE: 4
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213 1 5 10 15
215 Leu Asp Ala Cys
216 20
219 <210> SEQ ID NO: 5
220 <211> LENGTH: 9
221 <212> TYPE: PRT
222 <213> ORGANISM: Artificial Sequence
224 <220> FEATURE:
225 <223> OTHER INFORMATION: Description of Artificial Sequence: A peptidic
226 membrane binding element of SEQ ID NO: 4
228 <400> SEQUENCE: 5
229 Pro Ser Lys Lys Lys Lys Lys Lys Pro
230 1 5
233 <210> SEQ ID NO: 6
234 <211> LENGTH: 7
235 <212> TYPE: PRT
236 <213> ORGANISM: Artificial Sequence
238 <220> FEATURE:
239 <223> OTHER INFORMATION: Description of Artificial Sequence: A peptidic
240 membrane binding element of SEQ ID NO: 4
242 <400> SEQUENCE: 6
243 Leu Ser Ser Arg Leu Asp Ala
244 1 5
247 <210> SEQ ID NO: 7
248 <211> LENGTH: 16
249 <212> TYPE: PRT
250 <213> ORGANISM: Artificial Sequence
252 <220> FEATURE:
253 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of
254 electrostatic switch sequence
256 <400> SEQUENCE: 7
257 Asp Gly Pro Lys Lys Lys Lys Lys Lys Ser Pro Ser Lys Ser Ser Gly
258 1 5 10 15
261 <210> SEQ ID NO: 8
262 <211> LENGTH: 16
263 <212> TYPE: PRT
264 <213> ORGANISM: Artificial Sequence
266 <220> FEATURE:
267 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of
268 electrostatic switch sequence
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271 Gly Ser Ser Lys Ser Pro Ser Lys Lys Lys Lys Lys Lys Pro Gly Asp
272 1 5 10 15
275 <210> SEQ ID NO: 9

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276 <211> LENGTH: 20
277 <212> TYPE: PRT
278 <213> ORGANISM: Artificial Sequence
280 <220> FEATURE:
281 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of
282 electrostatic switch sequence
284 <400> SEQUENCE: 9
285 Ser Pro Ser Asn Glu Thr Pro Lys Lys Lys Lys Lys Arg Phe Ser Phe
286 1 5 10 15
288 Lys Lys Ser Gly
289 20
292 <210> SEQ ID NO: 10
293 <211> LENGTH: 16
294 <212> TYPE: PRT
295 <213> ORGANISM: Artificial Sequence
297 <220> FEATURE:
298 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of
299 electrostatic switch sequence
301 <400> SEQUENCE: 10
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303 1 5 10 15
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307 <211> LENGTH: 14
308 <212> TYPE: PRT
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311 <220> FEATURE:
312 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of
313 electrostatic switch sequence
315 <400> SEQUENCE: 11
316 Ser Lys Asp Gly Lys Lys Lys Lys Lys Ser Lys Thr Lys
317 1 5 10

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/936,205

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Input Set : A:\37945-0024.txt

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L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date